

SEQUENCE LISTING

<110> Oakley, Robert H.
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- <120> Methods of Screening Compositions for G Protein-Coupled Receptor Desensitization Inhibitory Activity
- <130> 033072-044
- <140> US 10/633,438
- <141> 2003-08-01
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- <151> 2001-11-05
- <150> US 60/245,772
- <151> 2000-11-03
- <150> US 60/260,363
- <151> 2001-01-08
- <160> 65
- <170> FastSEQ for Windows Version 4.0
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- <213> Homo sapiens
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Leu Lys Ile Trp Asn Asp His Phe Arg Cys Gln Pro Ala Pro Pro Ile 20 25 30

Asp Glu Asp Leu Pro Glu Glu Arg Pro Asp Asp 35. 40

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Asn Pro Ile Ile Tyr Pro Cys Ser Ser Lys Glu Phe Lys Arg Ala Phe 1 5 10 15

Val Arg Ile Leu Gly Cys Gln Cys Arg Gly Arg Gly Arg Arg Arg Arg 20 25 30

Arg Arg Arg Arg Leu Gly Gly Cys Ala Tyr Thr Tyr Arg Pro Trp

Thr Arg Gly Gly Ser Leu Glu Arg Ser Gln Ser Arg Lys Asp Ser Leu 50 55 60

Asp Asp Ser Gly Ser Cys Leu Ser Gly Ser Gln Arg Thr Leu Pro Ser 65 70 . 75 80

033072-044.ST25

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Ala Ser Pro Ser Pro Gly Tyr Leu Gly Arg Gly Ala Pro Pro Pro Val
Glu Leu Cys Ala Phe Pro Glu Trp Lys Ala Pro Gly Ala Leu Leu Ser
                                105
Leu Pro Ala Pro Glu Pro Pro Gly Arg Arg Gly Arg His Asp Ser Gly
                            120
Pro Leu Phe Thr Phe Lys Leu Leu Thr Glu Pro Glu Ser Pro Gly Thr
                        135
Asp Gly Gly Ala Ser Asn Gly Gly Cys Glu Ala Ala Ala Asp Val Ala
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Asn Gly Gln Pro Gly Phe Lys Ser Asn Met Pro Leu Ala Pro Gly Gln
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Phe
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Lys Lys Ile Leu Cys Arg Gly Asp Arg Lys Arg Ile Val
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Lys His Ile Leu Phe Arg Arg Arg Arg Gly Phe Arg Gln
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Asn Pro Ile Ile Tyr Cys Arg Ser Pro Asp Phe Arg Lys Ala Phe Gln
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Gly Leu Leu Cys Cys Ala Arg Arg Ala Ala Arg Arg Arg His Ala Thr
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His Gly Asp Arg Pro Arg Ala Ser Gly Cys Leu Ala Arg Pro Gly Pro
Pro Pro Ser Pro Gly Ala Ala Ser Asp Asp Asp Asp Asp Val Val
Gly Ala Thr Pro Pro Ala Arg Leu Leu Glu Pro Trp Ala Gly Cys Asn
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Gly Gly Ala Ala Ala Asp Ser Asp Ser Ser Leu Asp Glu Pro Cys Arg
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Pro Gly Phe Ala Ser Glu Ser Lys Val
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Glu Leu Leu Cys Leu Arg Arg Ser Ser Leu Lys Ala Tyr Gly Asn Gly
Tyr Ser Ser Asn Gly Asn Thr Gly Glu Gln Ser Gly Tyr His Val Glu
Gln Glu Lys Glu Asn Lys Leu Leu Cys Glu Asp Leu Pro Gly Thr Glu
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Gln Gly Arg Asn Cys Ser Thr Asn Asp Ser Leu Leu
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Thr Leu Leu Gly Cys Tyr Arg Leu Cys Pro Ala Thr Asn Asn Ala Ile
Glu Thr Val Ser Ile Asn Asn Asn Gly Ala Ala Met Phe Ser Ser His
                            40
His Glu Pro Arg Gly Ser Ile Ser Lys Glu Cys Asn Leu Val Tyr Leu
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Ile Pro His Ala Val Gly Ser Ser Glu Asp Leu Lys Lys Glu Glu Ala
                                        75
Ala Gly Ile Ala Arg Pro Leu Glu Lys Leu Ser Pro Ala Leu Ser Val
                85
Ile Leu Asp Tyr Asp Thr Asp Val Ser Leu Glu Lys Ile Gln Pro Ile
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                               105
Thr Gln Asn Gly Gln His Pro Thr
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Asn Pro Ile Ile Tyr Thr Thr Phe Asn Ile Glu Phe Arg Lys Ala Phe
                                    10
Leu Lys Ile Leu His Cys
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Asn Pro Val Ile Tyr Thr Thr Phe Asn Ile Glu Phe Arg Lys Ala Phe
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Leu Lys Ile Leu Ser Cys
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Arg Lys Ala Leu Arg Ala Cys Cys
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Gln Leu Leu Gly Cys Ser His Phe Cys Ser Arg Thr Pro Val Glu Thr
Val Asn Ile Ser Asn Glu Leu Ile Ser Tyr Asn Gln Asp Ile Val Phe
His Lys Glu Ile Ala Ala Tyr Ile His Met Met Pro Asn Ala Val
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Thr Pro Gly Asn Arg Glu Val Asp Asn. Asp Glu Glu Glu Gly Pro Phe
Asp Arg Met Phe Gln Ile Tyr Gln Thr Ser Pro Asp Gly Asp Pro Val
                                     90
                85
Ala Glu Ser Val Trp Glu Leu Asp Cys Glu Gly Glu Ile Ser Leu Asp
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Lys Ile Thr Pro Phe Thr Pro Asn Gly Phe His
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Pro Lys Arg Pro Gly Ser Val His Arg Thr Pro Ser Arg Gln Cys
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Lys His Leu Leu Met Cys His Tyr Lys Asn Ile Gly Ala Thr Arg
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Asn Pro Val Cys Tyr Ala Leu Cys Asn Lys Thr Phe Arg Thr Thr Lys
Met Leu Leu Cys Gln Cys Asp Lys Lys Lys Arg Arg Lys Gln Gln
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Tyr Gln Gln Arg Gln Ser Val Ile Phe His Lys Arg Ala Pro Glu Gln
Ala Leu
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Asn Pro Ala Cys Tyr Ala Leu Cys Asn Ala Thr Phe Lys Lys Thr Phe
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Arg His Leu Leu Cys Gln Tyr Arg Asn Ile Gly Thr Ala Arg
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<213> Homo sapiens

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Asn Pro Ile Cys Tyr Ala Leu Cys Asn Arg Thr Phe Arg Lys Thr Phe
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Lys Met Leu Leu Cys Arg Trp Lys Lys Lys Val Glu Glu Lys
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Leu Tyr Trp Gln Gly Asn Ser Lys Leu Pro
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Asn Pro Val Ile Tyr Ala Tyr Phe Asn Lys Asp Phe Gln Asn Ala Phe
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Lys Lys Ile Ile Lys Cys Lys Phe
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<211> 26
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Asn Pro Ile Ile Tyr Thr Met Ser Asn Glu Asp Phe Lys Gln Ala Phe
His Lys Leu Ile Arg Phe Lys Cys Thr Ser
<210> 20
<211> 24
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Asn Pro Leu Leu Tyr Thr Ser Phe Asn Glu Asp Phe Lys Leu Ala Phe
Lys Lys Leu Ile Arg Cys Arg Glu
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<210> 21
<211> 37
<212> PRT
<213> Homo sapiens
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Asn Pro Ile Ile Tyr Cys Leu Arg Asn Gln Glu Val Lys Arg Ala Leu
                                   10
Cys Cys Ile Leu His Leu Tyr Gln His Gln Asp Pro Asp Pro Lys Lys
                              25
           20
Gly Ser Arg Asn Val
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Arg Arg Leu Leu Gly Lys Gly Arg Glu Val Gly
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Asn Pro Leu Phe Tyr Gly Phe Leu Gly Lys Lys Phe Lys Arg Tyr Phe
Leu Gln Leu Leu Lys Tyr Ile Pro Pro Lys Ala Lys Ser His Ser Asn
                                 25
Leu Ser Thr Lys Met Ser Thr Leu Ser Tyr Arg Pro Ser Asp Asn Val
                            40
Ser Ser Ser Thr Lys Lys Pro Ala Pro Cys Phe Glu Val Glu
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Asn Pro Phe Leu Tyr Cys Phe Val Gly Asn Arg Phe Gln Gln Lys Leu
Arg Ser Val Phe Arg Val Pro Ile Thr Trp Leu Gln Gly Lys Arg Glu
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Ser Met Ser Cys Arg Lys Ser Ser Ser Leu Arg Glu Met Glu Thr Phe
Val Ser
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Asn Pro Leu Ile Tyr Ala Phe Ile Gly Gln Lys Phe Arg His Gly Leu
Leu Lys Ile Leu Ala Ile His Gly Leu Ser Lys Asp Ser Leu Pro Lys
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            20
Asp Ser Arg Pro Ser Phe Val Gly Ser Ser Ser Gly His Thr Ser Thr
Thr Leu
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Asn Pro Leu Ile Tyr Ala Phe Ala Gly Glu Lys Phe Arg Arg Tyr Leu
Tyr His Leu Tyr Gly Lys Cys Leu Ala Val Leu Cys Gly Arg Ser Val
                                25
His Val Asp Phe Ser Ser Glu Ser Gln Arg Ser Arg His Gly Ser
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Val Leu Ser Ser Asn Phe Thr Tyr His Thr Ser Asp Gly Asp Ala Leu
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Leu Leu Leu
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Leu Ala Thr Leu Ala Cys Leu Cys Pro Val Trp Arg Arg Arg Lys
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Arg Pro Ala Phe Ser Arg Lys Ala Asp Ser Val Ser Ser Asn His Thr
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Leu Ser Ser Asn Ala Thr Arg Glu Thr Leu Tyr
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Asn Pro Ile Ile Tyr Cys Cys Leu Asn Asp Arg Phe Arg Leu Gly Phe
Lys His Ala Phe Arg Cys Cys Pro Phe Ile Ser Ala Gly Asp Tyr Glu
                                25
Gly Leu Glu Met Lys Ser Thr Arg Tyr Leu Gln Thr Gln Gly Ser Val
                            40
Tyr Lys Val Ser Arg Leu Glu Thr Thr Ile Ser Thr Val Val Gly Ala
                        55
His Glu Glu Glu Pro Glu Asp Gly Pro Lys Ala Thr Pro Ser Ser Leu
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Asp Leu Thr Ser Asn Cys Ser Ser Arg Ser Asp Ser Lys Thr Met Thr
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Glu Ser Phe Ser Phe Ser Ser Asn Val Leu Ser
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<212> PRT

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Arg Ser Leu Leu Cys Cys Ala Arg Gly Arg Thr Pro Pro Ser Leu Gly
                            ' 25
Pro Gln Asp Glu Ser Cys Thr Thr Ala Ser Ser Ser Leu Ala Lys Asp
Thr Ser Ser
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Arg Lys Leu Cys Asn Cys Lys Gln Lys Pro Thr Glu Lys Pro Ala Asn
Tyr Ser Val Ala Leu Asn Tyr Ser Val Ile Lys Glu Ser Asp His Phe
                            40
Ser Thr Glu Leu Asp Asp Ile Thr Val Thr Asp Thr Tyr Leu Ser Ala
Thr Lys Val Ser Phe Asp Asp Thr Cys Leu Ala Ser Glu Val Ser Phe
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Ser Gln Ser
<210> 31
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Asn Pro Trp Ile Tyr Met Leu Phe Thr Gly His Leu Phe His Glu Leu
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Val Gln Arg Phe Leu Cys Cys Ser Ala Ser Tyr Leu Lys Gly Arg Arg
            20
                                25
Leu Gly Glu Thr Ser Ala Ser Lys Lys Ser Asn Ser Ser Ser Phe Val
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Leu Ser His Arg Ser Ser Ser Gln Arg Ser Cys Ser Gln Pro Ser Thr
Ala
65
<210> 32
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Asn Pro Val Leu Tyr Ser Leu Met Ser Ser Arg Phe Arg Glu Thr Phe
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Arg Ser Met Phe Pro Ser Cys Glu Gly Thr Ala Gln Pro Leu Asp Asn
Ser Met Gly Asp Ser Asp Cys Leu His Lys His Ala Asn Asn Ala Ala
        35
Ser Val His Arg Ala Ala Glu Ser Cys Ile Lys Ser Thr Val Lys Ile
Ala Lys Val Thr Met Ser Val Ser Thr Asp Thr Ser Ala Glu Ala Leu
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Asn Pro Val Leu Tyr Ala Phe Leu Asp Glu Asn Phe Lys Arg Cys Phe
Arg Gln Leu Cys Arg Lys Pro Cys Gly Arg Pro Asp Pro Ser Ser Phe
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Ser Arg Pro Arg Glu Ala Thr Ala Arg Glu Arg Val Thr Ala Cys Thr
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                            40
Pro Ser Asp Gly Pro Gly Gly Gly Arg Ala Ala
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Asp Pro Phe Val Tyr Tyr Phe Val Ser His Asp Phe Arg Asp His Ala
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Lys Asn Ala Leu Leu Cys Arg Ser Val Arg Thr Val Lys Gln Met Gln
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Val Ser Leu Thr Ser Lys Lys His Ser Arg Lys Ser Ser Ser Tyr Ser
                            40
Ser Ser Ser Thr Thr Val Lys Thr Ser Tyr
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Asn Gly Glu Val Gln Ala Glu Leu Arg Arg Lys Trp Arg Arg Trp His
Leu Gln Gly Val Leu Gly Trp Ser Ser Lys Ser Gln His Pro Trp Gly
Gly Ser Asn Gly Ala Thr Cys Ser Thr Gln Val Ser Met Leu Thr Arg
                           40
Val Ser Pro Ser Ala Arg Arg Ser Ser Ser Phe Gln Ala Glu Val Ser
Leu Val
65
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Met Leu Met Ala Ser Thr Thr Ser Ala Val Pro Gly His Pro Ser Leu
                5
                                   10
Pro Ser Leu Pro Ser Asn Ser Ser Gln Glu Arg Pro Leu Asp Thr Arg
                               25
Asp Pro Leu Leu Ala Arg Ala Glu Leu Ala Leu Leu Ser Ile Val Phe
                           40
Val Ala Val Ala Leu Ser Asn Gly Leu Val Leu Ala Ala Leu Ala Arg
                        55
Arg Gly Arg Arg Gly His Trp Ala Pro Ile His Val Phe Ile Gly His
                    70
                                       75
Leu Cys Leu Ala Asp Leu Ala Val Ala Leu Phe Gln Val Leu Pro Gln
               85
                                   90
Leu Ala Trp Lys Ala Thr Asp Arg Phe Arg Gly Pro Asp Ala Leu Cys
                               105
Arg Ala Val Lys Tyr Leu Cln Mct Val Gly Met Tyr Ala Ser Ser Tyr
       115
                           120
Met Ile Leu Ala Met Thr Leu Asp Arg His Arg Ala Ile Cys Arg Pro
                       135
Met Leu Ala Tyr Arg His Gly Ser Gly Ala His Trp Asn Arg Pro Val
                   150
                                       155
Leu Val Ala Trp Ala Phe Ser Leu Leu Ser Leu Pro Gln Leu Phe
               165
                                   170
Ile Phe Ala Gln Arg Asn Val Glu Gly Gly Ser Gly Val Thr Asp Cys
                               185
          180
Trp Ala Cys Phe Ala Glu Pro Trp Gly Arg Arg Thr Tyr Val Thr Trp
 . 195
                           200
Ile Ala Leu Met Val Phe Val Ala Pro Thr Leu Gly Ile Ala Ala Cys
                                           220
                       215
Gln Val Leu Ile Phe Arg Glu Ile His Ala Ser Leu Val Pro Gly Pro
                                       235
                   230
Ser Glu Arg Pro Gly Gly Arg Arg Gly Arg Arg Thr Gly Ser Pro
               245
                                   250
Gly Glu Gly Ala His Val Ser Ala Ala Val Ala Lys Thr Val Arg Met
                               265
           260
Thr Leu Val Ile Val Val Val Tyr Val Leu Cys Trp Ala Pro Phe Phe
                           280
Leu Val Gln Leu Trp Ala Ala Trp Asp Pro Glu Ala Pro Leu Glu Gly
                                        . 300
                       295
Ala Pro Phe Val Leu Leu Met Leu Leu Ala Ser Leu Asn Ser Cys Thr
                                        315
                    310
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Asn Pro Trp Ile Tyr Ala Ser Phe Ser Ser Ser Val Ser Ser Glu Leu
Arg Ser Leu Leu Cys Cys Ala Arg Gly Arg Thr Pro Pro Ser Leu Gly
                               345
Pro Gln Asp Glu Ser Cys Thr Thr Ala Ser Ser Leu Ala Lys Asp
                            360
Thr Ser Ser
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<213> Golden hamster
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Trp Gly Glu Leu Lys Asp Ala Asn Phe Thr Gly Pro Asn Gln Thr Ser
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Ser Asn Ser Thr Leu Pro Gln Leu Asp Val Thr Arg Ala Ile Ser Val
                            4.0
Gly Leu Val Leu Gly Ala Phe Ile Leu Phe Ala Ile Val Gly Asn Ile
                        55
Leu Val Ile Leu Ser Val Ala Cys Asn Arg His Leu Arg Thr Pro Thr
                    70
                                        75
Asn Tyr Phe Ile Val Asn Leu Ala Ile Ala Asp Leu Leu Ser Phe
                                   90
               85
Thr Val Leu Pro Phe Ser Ala Thr Leu Glu Val Leu Gly Tyr Trp Val
            100
                                105
Leu Gly Arg Ile Phe Cys Asp Ile Trp Ala Ala Val Asp Val Leu Cys
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                            120
Cys Thr Ala Ser Ile Leu Ser Leu Cys Ala Ile Ser Ile Asp Glu Tyr
                        135
                                            140
Ile Gly Val Arg Tyr Ser Leu Gln Tyr Pro Thr Leu Val Thr Arg Arg
                                       155
                    150
Lys Ala Ile Leu Ala Leu Leu Ser Val Trp Val Leu Ser Thr Val Ile
                165
                                    170
Ser Ile Gly Pro Leu Leu Gly Trp Lys Glu Pro Ala Pro Asn Asp Asp
                                185
Lys Glu Cys Gly Val Thr Glu Glu Pro Phe Tyr Ala Leu Phe Ser Ser
                            200
Leu Gly Ser Phe Tyr Ile Pro Leu Ala Val Ile Leu Val Met Tyr Cys
                        215
                                            220
Arg Val Tyr Ile Val Ala Lys Arg Thr Thr Lys Asn Leu Glu Ala Gly
                    230
                                        235
Val Met Lys Glu Met Ser Asn Ser Lys Glu Leu Thr Leu Arg Ile His
                245
                                    250
Ser Lys Asn Phe His Glu Asp Thr Leu Ser Ser Thr Lys Ala Lys Gly
                                265
                                                    270
His Asn Pro Arg Ser Ser Ile Ala Val Lys Leu Phe Lys Phe Ser Arg
                            280
        275
Glu Lys Lys Ala Ala Lys Thr Leu Gly Ile Val Val Gly Met Phe Ile
                        295
                                            300
Leu Cys Trp Leu Pro Phe Phe Ile Ala Leu Pro Leu Gly Ser Leu Phe
                                        315
                    310
Ser Thr Leu Lys Pro Pro Asp Ala Val Phe Lys Val Val Phe Trp Leu
                                    330
                325
Gly Tyr Phe Asn Ser Cys Leu Asn Pro Ile Ile Tyr Pro Cys Ser Ser
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340
                               345
Lys Glu Phe Lys Arg Ala Phe Met Arg Ile Leu Gly Cys Gln Cys Arg
                          360
                                              365
Ser Gly Arg Arg Arg Arg Arg Arg Arg Leu Gly Ala Cys Ala Tyr
                       375
Thr Tyr Arg Pro Trp Thr Arg Gly Gly Ser Leu Glu Arg Ser Gln Ser
                   390
                                       395
Arg Lys Asp Ser Leu Asp Asp Ser Gly Ser Cys Met Ser Gly Ser Gln
                                   410
               405
Arg Thr Leu Pro Ser Ala Ser Pro Ser Pro Gly Tyr Leu Gly Arg Gly
          420
                               425
Ala Gln Pro Pro Leu Glu Leu Cys Ala Tyr Pro Glu Trp Lys Ser Gly
                          440
Ala Leu Leu Ser Leu Pro Glu Pro Pro Gly Arg Arg Gly Arg Leu Asp
                       455
                                           460
Ser Gly Pro Leu Phe Thr Phe Lys Leu Leu Gly Glu Pro Glu Ser Pro
                                       475
                470
Gly Thr Glu Gly Asp Ala Ser Asn Gly Gly Cys Asp Ala Thr Thr Asp
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Leu Ala Asn Gly Gln Pro Gly Phe Lys Ser Asn Met Pro Leu Ala Pro
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Gly His Phe
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Trp Gly Glu Leu Lys Asp Ala Asn Phe Thr Gly Pro Asn Gln Thr Ser
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Ser Asn Ser Thr Leu Pro Gln Leu Asp Val Thr Arg Ala Ile Ser Val
Gly Leu Val Leu Gly Ala Phe Ile Leu Phe Ala Ile Val Gly Asn Ile
                       55
Leu Val Ile Leu Ser Val Ala Cys Asn Arg His Leu Arg Thr Pro Thr
                                       75
Asn Tyr Phe Ile Val Asn Leu Ala Ile Ala Asp Leu Leu Ser Phe
               85
                                   90
Thr Val Leu Pro Phe Ser Ala Thr Leu Glu Val Leu Gly Tyr Trp Val
                               105
Leu Gly Arg Ile Phe Cys Asp Ile Trp Ala Ala Val Asp Val Leu Cys
                           120
                                               125
       115
Cys Thr Ala Ser Ile Leu Ser Leu Cys Ala Ile Ser Ile Asp Ala Tyr
                       135
                                           140
Ile Gly Val Arg Tyr Ser Leu Gln Tyr Pro Thr Leu Val Thr Arg Arg
                   150
                                       155
Lys Ala Ile Leu Ala Leu Leu Ser Val Trp Val Leu Ser Thr Val Ile
                                   170
               165
Ser Ile Gly Pro Leu Leu Gly Trp Lys Glu Pro Ala Pro Asn Asp Asp
                               185
                                                   190
           180
Lys Glu Cys Gly Val Thr Glu Glu Pro Phe Tyr Ala Leu Phe Ser Ser
                           200
       195
Leu Gly Ser Phe Tyr Ile Pro Leu Ala Val Ile Leu Val Met Tyr Cys
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Arg Val Tyr Ile Val Ala Lys Arg Thr Thr Lys Asn Leu Glu Ala Gly

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Val Met Lys Glu Met Ser Asn Ser Lys Glu Leu Thr Leu Arg Ile His
               245
                                  250
Ser Lys Asn Phe His Glu Asp Thr Leu Ser Ser Thr Lys Ala Lys Gly
                               265
His Asn Pro Arg Ser Ser Ile Ala Val Lys Leu Phe Lys Phe Ser Arg
                           280
                                              285
Glu Lys Lys Ala Ala Lys Thr Leu Gly Ile Val Val Gly Met Phe Ile
                       295
                                          300
Leu Cys Trp Leu Pro Phe Phe Ile Ala Leu Pro Leu Gly Ser Leu Phe
                   310
                                       315
Ser Thr Leu Lys Pro Pro Asp Ala Val Phe Lys Val Val Phe Trp Leu
                                   330
               325
Gly Tyr Phe Asn Ser Cys Leu Asn Pro Ile Ile Tyr Pro Cys Ser Ser
                               345
Lys Glu Phe Lys Arg Ala Phe Met Arg Ile Leu Gly Cys Gln Cys Arg
                           360
Ser Gly Arg Arg Arg Arg Arg Arg Leu Gly Ala Cys Ala Tyr
                       375
Thr Tyr Arg Pro Trp Thr Arg Gly Gly Ser Leu Glu Arg Ser Gln Ser
                   390
Arg Lys Asp Ser Leu Asp Asp Ser Gly Ser Cys Met Ser Gly Ser Gln
                                   410
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Arg Thr Leu Pro Ser Ala Ser Pro Ser Pro Gly Tyr Leu Gly Arg Gly
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Ala Gln Pro Pro Leu Glu Leu Cys Ala Tyr Pro Glu Trp Lys Ser Gly
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Ala Leu Leu Ser Leu Pro Glu Pro Pro Gly Arg Arg Gly Arg Leu Asp
                       455
Ser Gly Pro Leu Phe Thr Phe Lys Leu Leu Gly Glu Pro Glu Ser Pro
                                       475
                   470
Gly Thr Glu Gly Asp Ala Ser Asn Gly Gly Cys Asp Ala Thr Thr Asp
                                  490
               485
Leu Ala Asn Gly Gln Pro Gly Phe Lys Ser Asn Met Pro Leu Ala Pro
Gly His Phe
       515
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Met Asn Pro Asp Leu Asp Thr Gly His Asn Thr Ser Ala Pro Ala Gln
Trp Gly Glu Leu Lys Asp Ala Asn Phe Thr Gly Pro Asn Gln Thr Ser
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                            40
Gly Leu Val Leu Gly Ala Phe Ile Leu Phe Ala Ile Val Gly Asn Ile
                        55
Leu Val Ile Leu Ser Val Ala Cys Asn Arg His Leu Arg Thr Pro Thr
                                        75
                    70
Asn Tyr Phe Ile Val Asn Leu Ala Ile Ala Asp Leu Leu Ser Phe
                85
                                   90
Thr Val Leu Pro Phe Ser Ala Thr Leu Glu Val Leu Gly Tyr Trp Val
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100
                              105
Leu Gly Arg Ile Phe Cys Asp Ile Trp Ala Ala Val Asp Val Leu Cys
                    120
Cys Thr Ala Ser Ile Leu Ser Leu Cys Ala Ile Ser Ile Asp His Tyr
                      135
Ile Gly Val Arg Tyr Ser Leu Gln Tyr Pro Thr Leu Val Thr Arg Arg
                  150
                                      155
Lys Ala Ile Leu Ala Leu Leu Ser Val Trp Val Leu Ser Thr Val Ile
                                  170
              165
Ser Ile Gly Pro Leu Leu Gly Trp Lys Glu Pro Ala Pro Asn Asp Asp
           180
                              185
Lys Glu Cys Gly Val Thr Glu Glu Pro Phe Tyr Ala Leu Phe Ser Ser
       195
                          200
Leu Gly Ser Phe Tyr Ile Pro Leu Ala Val Ile Leu Val Met Tyr Cys
                      215
Arg Val Tyr Ile Val Ala Lys Arg Thr Thr Lys Asn Leu Glu Ala Gly
                  230
                                     235
Val Met Lys Glu Met Ser Asn Ser Lys Glu Leu Thr Leu Arg Ile His
                                  250
               245
Ser Lys Asn Phe His Glu Asp Thr Leu Ser Ser Thr Lys Ala Lys Gly
           260
                              265
His Asn Pro Arg Ser Ser Ile Ala Val Lys Leu Phe Lys Phe Ser Arg
                           280
Glu Lys Lys Ala Ala Lys Thr Leu Gly Ile Val Val Gly Met Phe Ile
                      295
                                          300
Leu Cys Trp Leu Pro Phe Phe Ile Ala Leu Pro Leu Gly Ser Leu Phe
                   310
                                      315
Ser Thr Leu Lys Pro Pro Asp Ala Val Phe Lys Val Val Phe Trp Leu
                                   330
                                                      335
               325
Gly Tyr Phe Asn Ser Cys Leu Asn Pro Ile Ile Tyr Pro Cys Ser Ser
                               345
Lys Glu Phe Lys Arg Ala Phe Met Arg Ile Leu Gly Cys Gln Cys Arg
                           360
       355
Ser Gly Arg Arg Arg Arg Arg Arg Arg Lou Gly Ala Cys Ala Tyr
                       375
                                          380
Thr Tyr Arg Pro Trp Thr Arg Gly Gly Ser Leu Glu Arg Ser Gln Ser
                  390
                                      395
Arg Lys Asp Ser Leu Asp Asp Ser Gly Ser Cys Met Ser Gly Ser Gln
     . 405
                                  410
Arg Thr Leu Pro Ser Ala Ser Pro Ser Pro Gly Tyr Leu Gly Arg Gly
          420
                              425
                                                  430
Ala Gln Pro Pro Leu Glu Leu Cys Ala Tyr Pro Glu Trp Lys Ser Gly
                           440
                                              445
Ala Leu Leu Ser Leu Pro Glu Pro Pro Gly Arg Arg Gly Arg Leu Asp
                      455
                                          460
Ser Gly Pro Leu Phe Thr Phe Lys Leu Gly Glu Pro Glu Ser Pro
                                      475
                  470
Gly Thr Glu Gly Asp Ala Ser Asn Gly Gly Cys Asp Ala Thr Thr Asp
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                                  490
Leu Ala Asn Gly Gln Pro Gly Phe Lys Ser Asn Met Pro Leu Ala Pro
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Gly His Phe
       515
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<210> 44

<211> 515

<212> PRT

<213> Golden hamster

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Met Asn Pro Asp Leu Asp Thr Gly His Asn Thr Ser Ala Pro Ala Gln
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Trp Gly Glu Leu Lys Asp Ala Asn Phe Thr Gly Pro Asn Gln Thr Ser
                               25
Ser Asn Ser Thr Leu Pro Gln Leu Asp Val Thr Arg Ala Ile Ser Val
                          40
Gly Leu Val Leu Gly Ala Phe Ile Leu Phe Ala Ile Val Gly Asn Ile
                       55
Leu Val Ile Leu Ser Val Ala Cys Asn Arg His Leu Arg Thr Pro Thr
                   70
                                      75
Asn Tyr Phe Ile Val Asn Leu Ala Ile Ala Asp Leu Leu Ser Phe
                                   90
Thr Val Leu Pro Phe Ser Ala Thr Leu Glu Val Leu Gly Tyr Trp Val
                               105
Leu Gly Arg Ile Phe Cys Asp Ile Trp Ala Ala Val Asp Val Leu Cys
                          120
Cys Thr Ala Ser Ile Leu Ser Leu Cys Ala Ile Ser Ile Asp Asn Tyr
                       135
Ile Gly Val Arg Tyr Ser Leu Gln Tyr Pro Thr Leu Val Thr Arg Arg
                  150
Lys Ala Ile Leu Ala Leu Leu Ser Val Trp Val Leu Ser Thr Val Ile
               165
                                  170
Ser Ile Gly Pro Leu Leu Gly Trp Lys Glu Pro Ala Pro Asn Asp Asp
                              185
Lys Glu Cys Gly Val Thr Glu Glu Pro Phe Tyr Ala Leu Phe Ser Ser
                          200
Leu Gly Ser Phe Tyr Ile Pro Leu Ala Val Ile Leu Val Met Tyr Cys
                       215
Arg Val Tyr Ile Val Ala Lys Arg Thr Thr Lys Asn Leu Glu Ala Gly
                   230
                                      235
Val Met Lys Glu Met Ser Asn Ser Lys Glu Leu Thr Leu Arg Ile His
                                  250
               245
Ser Lys Asn Phe His Glu Asp Thr Leu Ser Ser Thr Lys Ala Lys Gly
                               265
           260
His Asn Pro Arg Ser Ser Ile Ala Val Lys Leu Phe Lys Phe Ser Arg
                           280
Glu Lys Lys Ala Ala Lys Thr Leu Gly Ile Val Val Gly Met Phe Ile
                       295
Leu Cys Trp Leu Pro Phe Phe Ile Ala Leu Pro Leu Gly Ser Leu Phe
                   310
                                       315
Ser Thr Leu Lys Pro Pro Asp Ala Val Phe Lys Val Val Phe Trp Leu
               325
                                   330
Gly Tyr Phe Asn Ser Cys Leu Asn Pro Ile Ile Tyr Pro Cys Ser Ser
           340
                               345
Lys Glu Phe Lys Arg Ala Phe Met Arg Ile Leu Gly Cys Gln Cys Arg
                           360
       355
Ser Gly Arg Arg Arg Arg Arg Arg Arg Leu Gly Ala Cys Ala Tyr
                                          380
                       375
Thr Tyr Arg Pro Trp Thr Arg Gly Gly Ser Leu Glu Arg Ser Gln Ser
                   390
                                       395
Arg Lys Asp Ser Leu Asp Asp Ser Gly Ser Cys Met Ser Gly Ser Gln
                                   410
               405
Arg Thr Leu Pro Ser Ala Ser Pro Ser Pro Gly Tyr Leu Gly Arg Gly
                               425
           420
Ala Gln Pro Pro Leu Glu Leu Cys Ala Tyr Pro Glu Trp Lys Ser Gly
                           440
Ala Leu Leu Ser Leu Pro Glu Pro Pro Gly Arg Arg Gly Arg Leu Asp
                       455
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      Ser Gly Pro Leu Phe Thr Phe Lys
      Leu Leu Gly Glu Pro Glu Ser Pro 465
      470
      475
      480

      Gly Thr Glu Gly Asp Ala Ser Asn Gly Gly Cys
      Gly Gly Cys Asp Ala Thr Thr Asp 490
      495

      Leu Ala Asn Gly Gln Pro Gly Phe Lys Ser Asn Met Pro Leu Ala Pro 505
      500

      Gly His Phe 515
      515
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<210> 45 <211> 359 <212> PRT <213> Rattus norvegicus

<400> 45

Met Ala Leu Asn Ser Ser Ala Glu Asp Gly Ile Lys Arg Ile Gln Asp Asp Cys Pro Lys Ala Gly Arg His Ser Tyr Ile Phe Val Met Ile Pro Thr Leu Tyr Ser Ile Ile Phe Val Val Gly Ile Phe Gly Asn Ser Leu 40 Val Val Ile Val Ile Tyr Phe Tyr Met Lys Leu Lys Thr Val Ala Ser 55 Val Phe Leu Leu Asn Leu Ala Leu Ala Asp Leu Cys Phe Leu Leu Thr Cys Pro Leu Trp Ala Val Tyr Thr Ala Met Glu Tyr Arg Trp Pro Phe 90 Gly Asn His Leu Cys Lys Ile Ala Ser Ala Ser Val Thr Phe Asn Leu 105 Tyr Ala Ser Val Phe Leu Leu Thr Cys Leu Ser Ile Asp Arg Tyr Leu 120 Ala Ile Val His Pro Met Lys Ser Arg Leu Arg Arg Thr Met Leu Val 135 Ala Lys Val Thr Cys Ile Ile Ile Trp Leu Met Ala Gly Leu Ala Ser 150 155 Leu Pro Ala Val Ile His Arg Asn Val Tyr Phe Ile Glu Asn Thr Asn 170 165 Ile Thr Val Cys Ala Phe His Tyr Glu Ser Arg Asn Ser Thr Leu Pro 185 Ile Gly Leu Gly Leu Thr Lys Asn Ile Leu Gly Phe Leu Phe Pro Phe 200 Leu Ile Ile Leu Thr Ser Tyr Thr Leu Ile Trp Lys Ala Leu Lys Lys 215 220 Ala Tyr Glu Ile Gln Lys Asn Lys Pro Arg Asn Asp Asp Ile Phe Arg 230 235 Ile Ile Met Ala Ile Val Leu Phe Phe Phe Phe Ser Trp Val Pro His 245 250 Gln Ile Phe Thr Phe Leu Asp Val Leu Ile Gln Leu Gly Val Ile His 265 260 Asp Cys Lys Ile Ser Asp Ile Val Asp Thr Ala Met Pro Ile Thr Ile 275 280 Cys Ile Ala Tyr Phe Asn Asn Cys Leu Asn Pro Leu Phe Tyr Gly Phe 295 300 Leu Gly Lys Lys Phe Lys Lys Tyr Phe Leu Gln Leu Leu Lys Tyr Ile 310 315 Pro Pro Lys Ala Lys Ser His Ser Ser Leu Ser Thr Lys Met Ser Thr 325 330 Leu Ser Tyr Arg Pro Ser Asp Asn Met Ser Ser Ser Ala Lys Lys Pro

345

340

Ala Ser Cys Phe Glu Val Glu 355 <210> 46 <211> 346 <212> PRT <213> Homo sapiens <400> 46 Met Met Trp Gly Ala Gly Ser Pro Leu Ala Trp Leu Ser Ala Gly Ser 10 Gly Asn Val Asn Val Ser Ser Val Gly Pro Ala Glu Gly Pro Thr Gly 25 Pro Ala Ala Pro Leu Pro Ser Pro Lys Ala Trp Asp Val Val Leu Cys 40 Ile Ser Gly Thr Leu Val Ser Cys Glu Asn Ala Leu Val Val Ala Ile Ile Val Gly Thr Pro Ala Phe Arg Ala Pro Met Phe Leu Leu Val Gly Ser Leu Ala Val Ala Asp Leu Leu Ala Gly Leu Gly Leu Val Leu His 90 Phe Ala Ala Val Phe Cys Ile Gly Ser Ala Glu Met Ser Leu Val Leu 105 Val Gly Val Leu Ala Met Ala Phe Thr Ala Ser Ile Gly Ser Leu Leu 120 Ala Ile Thr Val Asp Arg Tyr Leu Ser Leu Tyr Asn Ala Leu Thr Tyr 135 Tyr Ser Glu Thr Thr Val Thr Arg Thr Tyr Val Met Leu Ala Leu Val 150 155 Trp Gly Gly Ala Leu Gly Leu Gly Leu Pro Val Leu Ala Trp Asn 165 170 Cys Leu Asp Gly Leu Thr Thr Cys Gly Val Val Tyr Pro Leu Ser Lys 180 185 Asn His Leu Val Val Leu Ala Ile Ala Phe Phe Met Val Phe Gly Ile 200 Met Leu Gln Leu Tyr Ala Gln Ile Cys Arg Ile Val Cys Arg His Ala 215 Gln Gln Ile Ala Leu Gln Arg His Leu Leu Pro Ala Ser His Tyr Val 235 Ala Thr Arg Lys Gly Ile Ala Thr Leu Ala Val Val Leu Gly Ala Phe 245 Ala Ala Cys Trp Leu Pro Phe Thr Val Tyr Cys Leu Leu Gly Asp Ala 260 265 His Ser Pro Pro Leu Tyr Thr Tyr Leu Thr Leu Leu Pro Ala Thr Tyr 280 Asn Ser Met Ile Asn Pro Ile Ile Tyr Ala Phe Arg Asn Gln Asp Val 295 Gln Lys Val Leu Trp Ala Val Cys Cys Cys Cys Ala Ala Ala Arg Gly 310 315 Arg Thr Pro Pro Ser Leu Gly Pro Gln Asp Glu Ser Cys Thr Thr Ala 325 Ser Ser Ser Leu Ala Lys Asp Thr Ser Ser

<210> 47 <211> 1041

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<212> DNA
<213> Homo sapiens
<400> 47
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gtaagcagcg tgggcccagc agaggggccc acaggtccag ccgcaccact gccctcgcct 120
aaggeetggg atgtggtget etgeatetea ggeaceetgg tgteetgega gaatgegeta 180
gtggtggcca tcatcgtggg cactcctgcc ttccgtgccc ccatgttcct gctggtgggc 240
agectggeeg tggeagacet getggeagge etgggeetgg teetgeactt tgetgetgte 300
ttctgcatcg gctcagcgga gatgagcctg gtgctggttg gcgtgctggc aatggccttt 360
acygocagca toggoagtot actggocato actgtogaco gotacottto totgtacaat 420
gccctcacct actattcaga gacaacagtg acacggacct atgtgatgct ggccttagtg 480
tggggaggtg ccctgggcct ggggctgctg cctgtgctgg cctggaactg cctggatggc 540
ctgaccacat gtggcgtggt ttatccactc tccaagaacc atctggtagt tctggccatt 600
gccttcttca tggtgtttgg catcatgctg cagctctacg cccaaatctg ccgcatcgtc 660
tgccgccatg cccagcagat tgcccttcag cggcacctgc tgcctgcctc ccactatgtg 720
gccacccgca agggcattgc cacactggcc gtggtgcttg gagcctttgc cgcctgctgg 780
ttgcccttca ctgtctactg cctgctgggt gatgcccact ctccacctct ctacacctat 840
cttaccttgc tecetgecae ctacaactee atgateaace etateateta egeetteege 900
aaccaggatg tgcagaaagt gctgtgggct gtctgctgct gctgtgcggc cgcacgggga 960
cgcaccccac ccagcctqqq tccccaaqat qaqtcctgca ccaccqccaq ctcctccctq 1020
qccaaqqaca cttcatcqtq a
<210> 48
<211> 378
<212> PRT
<213> Homo sapiens
<400> 48
Met Asn Ala Ser Ala Ala Ser Leu Asn Asp Ser Gln Val Val Val Val
Ala Ala Glu Gly Ala Ala Ala Ala Thr Ala Ala Gly Gly Pro Asp
Thr Gly Glu Trp Cly Pro Pro Ala Ala Ala Ala Leu Gly Ala Gly Gly
                            40
Gly Ala Asn Gly Ser Leu Glu Leu Ser Ser Gln Leu Ser Ala Gly Pro
                        55
Pro Gly Leu Leu Pro Ala Val Asn Pro Trp Asp Val Leu Leu Cys
Val Ser Gly Thr Val Ile Ala Gly Glu Asn Ala Leu Val Val Ala Leu
Ile Ala Ser Thr Pro Ala Leu Arg Thr Pro Met Phe Val Leu Val Gly
                                105
Ser Leu Ala Thr Ala Asp Leu Leu Ala Gly Cys Gly Leu Ile Leu His
                            120
Phe Val Phe Gln Tyr Leu Val Pro Ser Glu Thr Val Ser Leu Leu Thr
                        135
                                            140
Val Gly Phe Leu Val Ala Ser Phe Ala Ala Ser Val Ser Ser Leu Leu
                                        155
                    150
Ala Ile Thr Val Asp Arg Tyr Leu Ser Leu Tyr Asn Ala Leu Thr Tyr
                                    170
                165
Tyr Ser Arg Arg Thr Leu Leu Gly Val His Leu Leu Leu Ala Ala Thr
            180
                                185
Trp Thr Val Ser Leu Gly Leu Gly Leu Leu Pro Val Leu Gly Trp Asn
                            200
Cys Leu Ala Glu Arg Ala Ala Cys Ser Val Val Arg Pro Leu Ala Arg
                        215
                                            220
Ser His Val Ala Leu Leu Ser Ala Ala Phe Phe Met Val Phe Gly Ile
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Met Leu His Leu Tyr Val Arg Ile Cys Gln Val Val Trp Arg His Ala
                245
                                    250
His Gln Ile Ala Leu Gln Gln His Cys Leu Ala Pro Pro His Leu Ala
                                265
                                                    270
Ala Thr Arg Lys Gly Val Gly Thr Leu Ala Val Val Leu Gly Thr Phe
        275
                            280
                                                 285
Gly Ala Ser Trp Leu Pro Phe Ala Ile Tyr Cys Val Val Gly Ser His
                        295
                                            300
Glu Asp Pro Ala Val Tyr Thr Tyr Ala Thr Leu Leu Pro Ala Thr Tyr
                    310
                                        315
Asn Ser Met Ile Asn Pro Ile Ile Tyr Ala Phe Arg Asn Gln Glu Ile
                                    330
Gln Arg Ala Leu Trp Leu Leu Leu Cys Gly Cys Ala Ala Ala Arg Gly
                                345
Arg Thr Pro Pro Ser Leu Gly Pro Gln Asp Glu Ser Cys Thr Thr Ala
                            360
Ser Ser Ser Leu Ala Lys Asp Thr Ser Ser
                        375
<210> 49
<211> 1137
<212> DNA
<213> Homo sapiens
<400> 49
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geggeggegg eggeeacage ageagggggg eeggaeaegg gegaatgggg acceeetget 120
geggeggete taggageegg eggeggaget aatgggtete tggagetgte etegeagetg 180
teggetggge caeegggaet cetgetgeea geggtgaate egtgggaegt geteetgtge 240
gtgtcgggga cagtgatcgc tggagaaaac gcgctggtgg tggcgctcat cgcgtccact 300
ccggcgctgc gcacgcccat gttcgtgctg gtaggcagcc tggccaccgc tgacctgttg 360
qcqqqctqtq qcctcatctt qcactttqtq ttccaqtact tqqtqccctc qqaqactqtq 420
agtotgotoa oggtgggott octogtggoo toottogoog octotgtoag cagootgotg 480
gecattacgg tggaccgcta cctgtccctg tataacgcgc tcacctatta ctcgcgccqq 540
accetgttgg gegtgeacet cetgettgee gecaettgga eegtgteeet aggeetgggg 600
ctgctgcccg tgctgggctg gaactgcctg gcagagcgcg ccgcctgcag cgtggtgcgc 660
ccgctggcgc gcagccacgt ggctctgctc tccgccgcct tcttcatggt cttcggcatc 720
atgctgcacc tgtacgtgcg catctgccag gtggtctggc gccacgcgca ccagatcgcg 780
ctgcagcagc actgcctggc gccaccccat ctcgctgcca ccagaaaggg tgtgggtaca 840
ctggctgtgg tgctgggcac tttcggcgcc agctggctgc ccttcgccat ctattgcgtg 900
gtgggcagcc atgaggaccc ggcggtctac acttacgcca ccctgctgcc cgccacctac 960
aactccatqa tcaatcccat catctatgcc ttccgcaacc aggagatcca gcgcgccctg 1020
tggctcctgc tctgtggctg tgcggccgca cggggacgca ccccacccag cctgggtccc 1080
caagatgagt cctqcaccac cgccagctcc tccctggcca aggacacttc atcgtga
<210> 50
<211> 350
<212> PRT
<213> Homo sapiens
<400> 50
Met Asn Glu Asp Leu Lys Val Asn Leu Ser Gly Leu Pro Arg Asp Tyr
                                    10
Leu Asp Ala Ala Ala Glu Asn Ile Ser Ala Ala Val Ser Ser Arg
                                 25
Val Pro Ala Val Glu Pro Glu Pro Glu Leu Val Val Asn Pro Trp Asp
                            40
Ile Val Leu Cys Thr Ser Gly Thr Leu Ile Ser Cys Glu Asn Ala Ile
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Val Val Leu Ile Ile Phe His Asn Pro Ser Leu Arg Ala Pro Met Phe

50

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70
                                        75
Leu Leu Ile Gly Ser Leu Ala Leu Ala Asp Leu Leu Ala Gly Ile Gly
                                    90
Leu Ile Thr Asn Phe Val Phe Ala Tyr Leu Leu Gln Ser Glu Ala Thr
                                105
Lys Leu Val Thr Ile Gly Leu Ile Val Ala Ser Phe Ser Ala Ser Val
                            120
Cys Ser Leu Leu Ala Ile Thr Val Asp Arg Tyr Leu Ser Leu Tyr Tyr
                        135
Ala Leu Thr Tyr His Ser Glu Arg Thr Val Thr Phe Thr Tyr Val Met
                    150
                                        155
Leu Val Met Leu Trp Gly Thr Ser Ile Cys Leu Gly Leu Leu Pro Val
                                    170
                165
Met Gly Trp Asn Cys Leu Arg Asp Glu Ser Thr Cys Ser Val Val Arg
                                185
Pro Leu Thr Lys Asn Asn Ala Ala Ile Leu Ser Val Ser Phe Leu Phe
                            200
Met Phe Ala Leu Met Leu Gln Leu Tyr Ile Gln Ile Cys Lys Ile Val
                        215
Met Arg His Ala His Gln Ile Ala Leu Gln His His Phe Leu Ala Thr
                                        235
                    230
Ser His Tyr Val Thr Thr Arg Lys Gly Val Ser Thr Leu Ala Ile Ile
                                    250
                245
Leu Gly Thr Phe Ala Ala Cys Trp Met Pro Phe Thr Leu Tyr Ser Leu
                                265
                                                     270
            260
Ile Ala Asp Tyr Thr Tyr Pro Ser Ile Tyr Thr Tyr Ala Thr Leu Leu
                            280
Pro Ala Thr Tyr Asn Ser Ile Ile Asn Pro Val Ile Tyr Ala Phe Arg
                        295
                                             300
Asn Gln Glu Ile Gln Lys Ala Leu Cys Leu Ile Cys Cys Gly Cys Ala
                    310
                                        315
Ala Ala Arg Gly Arg Thr Pro Pro Ser Lou Gly Pro Gln Asp Glu Ser
                                    330
                325
Cys Thr Thr Ala Ser Ser Ser Leu Ala Lys Asp Thr Ser Ser
            340
                                345
<210> 51
<211> 1053
<212> DNA
<213> Homo sapiens
<400> 51
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getgeggaga acatetegge tgetgtetee teeegggtte etgeegtaga geeagageet 120
gagetegtag teaacecetg ggacattgte ttgtgtacet egggaacect cateteetgt 180
qaaaatqcca ttqtqqtcct tatcatcttc cacaacccca gcctgcgagc acccatgttc 240
ctgctaatag gcagcctggc tcttgcagac ctgctggccg gcattggact catcaccaat 300
tttgtttttg cctacctgct tcagtcagaa gccaccaagc tggtcacgat cggcctcatt 360
gtcgcctctt tctctgcctc tgtctgcagc ttgctggcta tcactgttga ccgctacctc 420
toactgtact acgctctgac gtaccattcg gagaggacgg tcacgtttac ctatgtcatg 480
ctcgtcatgc tctgggggac ctccatctgc ctggggctgc tgcccgtcat gggctggaac 540
tgcctccgag acgagtccac ctgcagcgtg gtcagaccgc tcaccaagaa caacgcggcc 600
atcetetegg tgteetteet etteatgttt gegeteatge tteageteta cateeagate 660
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tgtaagattg tgatgaggca cgcccatcag atagccctgc agcaccactt cctggccacg 720 tcgcactatg tgaccacccg gaaaggggtc tccaccctgg ctatcatcct ggggacgttt 780 qctqcttqct ggatgccttt caccctctat tccttgatag cggattacac ctaccctcc 840

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atctatacct acgccaccct cctgcccgcc acctacaatt ccatcatcaa ccctgtcata 900
tatgetttea gaaaceaaga gateeagaaa gegetetgte teatttgetg eggetgegeg 960
gccgcacggg gacgcacccc acccagcctg ggtccccaag atgagtcctg caccaccgcc 1020
agctcctccc tggccaagga cacttcatcg tga
<210> 52
<211> 388
<212> PRT
<213> Homo sapiens
<400> 52
Met Ala Asn Thr Thr Gly Glu Pro Glu Glu Val Ser Gly Ala Leu Ser
Pro Pro Ser Ala Ser Ala Tyr Val Lys Leu Val Leu Leu Gly Leu Ile
                                25
Met Cys Val Ser Leu Ala Gly Asn Ala Ile Leu Ser Leu Leu Val Leu
                            40
Lys Glu Arg Ala Leu His Lys Ala Pro Tyr Tyr Phe Leu Leu Asp Leu
                        55
Cys Leu Ala Asp Gly Ile Arg Ser Ala Val Cys Phe Pro Phe Val Leu
                    70
                                        75
Ala Ser Val Arg His Gly Ser Ser Trp Thr Phe Ser Ala Leu Ser Cys
                                    90
Lys Ile Val Ala Phe Met Ala Val Leu Phe Cys Phe His Ala Ala Phe
                                105
Met Leu Phe Cys Ile Ser Val Thr Arg Tyr Met Ala Ile Ala His His
                            120
Arg Phe Tyr Ala Lys Arg Met Thr Leu Trp Thr Cys Ala Ala Val Ile
                        135
                                            140
Cys Met Ala Trp Thr Leu Ser Val Ala Met Ala Phe Pro Pro Val Phe
                    150
Asp Val Gly Thr Tyr Lys Phe Ile Arg Glu Glu Asp Gln Cys Ile Phe
                165
                                    170
Glu His Arg Tyr Phe Lys Ala Asn Asp Thr Leu Gly Phe Met Leu Met
                                185
Leu Ala Val Leu Met Ala Ala Thr His Ala Val Tyr Gly Lys Leu Leu
                            200
                                                205
Leu Phe Glu Tyr Arg His Arg Lys Met Lys Pro Val Gln Met Val Pro
                        215
Ala Ile Ser Gln Asn Trp Thr Phe His Gly Pro Gly Ala Thr Gly Gln
                    230
                                        235
Ala Ala Asn Trp Ile Ala Gly Phe Gly Arg Gly Pro Met Pro Pro
                245
                                    250
Thr Leu Leu Gly Ile Arg Gln Asn Gly His Ala Ala Ser Arg Arg Leu
            260
                                265
Leu Gly Met Asp Glu Val Lys Gly Glu Lys Gln Leu Gly Arg Met Phe
        275
                            280
                                                285
Tyr Ala Ile Thr Leu Leu Phe Leu Leu Leu Trp Ser Pro Tyr Ile Val
                                            300
                        295
Ala Cys Tyr Trp Arg Val Phe Val Lys Ala Cys Ala Val Pro His Arg
                    310
                                        315
Tyr Leu Ala Thr Ala Val Trp Met Ser Phe Ala Gln Ala Ala Val Asn
                325
                                    330
Pro Ile Val Cys Phe Leu Leu Asn Lys Asp Leu Lys Lys Cys Leu Arg
                                345
            340
Thr His Ala Pro Cys Ala Ala Ala Arg Gly Arg Thr Pro Pro Ser Leu
                            360
                                                365
Gly Pro Gln Asp Glu Ser Cys Thr Thr Ala Ser Ser Ser Leu Ala Lys
                        375
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Asp Thr Ser Ser
385
<210> 53
<211> 1167
<212> DNA
<213> Homo sapiens
<400> 53
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gccatcttgt ccctgctggt gctcaaggag cgtgccctgc acaaggctcc ttactacttc 180
ctgctggacc tgtgcctggc cgatggcata cgctctgccg tctgcttccc ctttgtgctg 240
gettetgtge gecaeggete tteatggaee tteagtgeae teagetgeaa gattgtggee 300 -
tttatggccg tqctcttttg cttccatgcg gccttcatgc tgttctgcat cagcgtcacc 360
cgctacatgg ccatcgccca ccaccgcttc tacgccaagc gcatgacact ctggacatgc 420
geggetgtea tetgeatgge etggaecetg tetgtggeea tggeetteee acetgtettt 480
gacgtgggca cctacaagtt tattcgggag gaggaccagt gcatctttga gcatcgctac 540
ttcaaqqcca atqacacqct qqqcttcatq cttatqttqq ctqtqctcat qqcaqctacc 600
catgctqtct acqqcaaqct qctcctcttc qaqtatcqtc accqcaaqat qaaqccaqtq 660
cagatggtgc cagccatcag ccagaactgg acattccatg gtcccggggc caccggccag 720
qctqctqcca actqgatcqc cggctttgqc cgtgggccca tgccaccaac cctgctgggt 780
atccggcaga atgggcatgc agccagccgg cggctactgg gcatggacga ggtcaagggt 840
qaaaaqcaqc tqqqccqcat qttctacqcq atcacactqc tctttctqct cctctqqtca 900
ccctacatcg tggcctgcta ctggcgagtg tttgtgaaag cctgtgctgt gccccaccgc 960
tacctggcca ctgctgtttg gatgagette geecaggetg cegteaacce aattgtetge 1020
ttcctqctca acaaqqacct caaqaaqtqc ctqaqqactc acqccccctq cqcqqccqca 1080
eggggaegea ceceacecag cetgggteee caagatgagt cetgeaceae egeeagetee 1140
tccctggcca aggacacttc atcgtga
                                                                   1167
<210> 54
<211> 388
<212> PRT
<213> Homo sapiens
<400> 54
Met Ala Asn Tyr Ser His Ala Ala Asp Asn Ile Leu Gln Asn Leu Ser
Pro Leu Thr Ala Phe Leu Lys Leu Thr Ser Leu Gly Phe Ile Ile Gly
Val Ser Val Val Gly Asn Leu Leu Ile Ser Ile Leu Leu Val Lys Asp
Lys Thr Leu His Arg Ala Pro Tyr Tyr Phe Leu Leu Asp Leu Cys Cys
Ser Asp Ile Leu Arg Ser Ala Ile Cys Phe Pro Phe Val Phe Asn Ser
                    70
Val Lys Asn Gly Ser Thr Trp Thr Tyr Gly Thr Leu Thr Cys Lys Val
Ile Ala Phe Leu Gly Val Leu Ser Cys Phe His Thr Ala Phe Met Leu
                                105
            100
Phe Cys Ile Ser Val Thr Arg Tyr Leu Ala Ile Ala His His Arg Phe
                            120
        115
Tyr Thr Lys Arg Leu Thr Phe Trp Thr Cys Leu Ala Val Ile Cys Met
                        135
Val Trp Thr Leu Ser Val Ala Met Ala Phe Pro Pro Val Leu Asp Val
                    150
                                        155
Gly Thr Tyr Ser Phe Ile Arg Glu Glu Asp Gln Cys Thr Phe Gln His
                165
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Arg Ser Phe Arg Ala Asn Asp Ser Leu Gly Phe Met Leu Leu Ala
            180
                                185
Leu Ile Leu Leu Ala Thr Gln Leu Val Tyr Leu Lys Leu Ile Phe Phe
                            200
                                                205
Val His Asp Arg Arg Lys Met Lys Pro Val Gln' Phe Val Ala Ala Val
                        215
                                            220
Ser Gln Asn Trp Thr Phe His Gly Pro Gly Ala Ser Gly Gln Ala Ala
                    230
                                        235
Ala Asn Trp Leu Ala Gly Phe Gly Arg Gly Pro Thr Pro Pro Thr Leu
                245
                                    250
Leu Gly Ile Arg Gln Asn Ala Asn Thr Thr Gly Arg Arg Leu Leu
            260
                                265
                                                    270
Val Leu Asp Glu Phe Lys Met Glu Lys Arg Ile Ser Arg Met Phe Tyr
                            280
Ile Met Thr Phe Leu Phe Leu Thr Leu Trp Gly Pro Tyr Leu Val Ala
                        295
                                            300
Cys Tyr Trp Arg Val Phe Ala Arg Gly Pro Val Val Pro Gly Gly Phe
                    310
                                        315
Leu Thr Ala Ala Val Trp Met Ser Phe Ala Gln Ala Gly Ile Asn Pro
                325
                                    330
Phe Val Cys Ile Phe Ser Asn Arg Glu Leu Arg Arg Cys Phe Ser Thr
                                345
                                                     350
Thr Leu Leu Tyr Cys Ala Ala Ala Arg Gly Arg Thr Pro Pro Ser Leu
                            360
                                                365
Gly Pro Gln Asp Glu Ser Cys Thr Thr Ala Ser Ser Ser Leu Ala Lys
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Asp Thr Ser Ser
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<211> 1167
<212> DNA
<213> Homo sapiens
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tttctgaaac tgacttcctt gggtttcata ataggagtca gcgtggtggg caacctcctg 120
atctccattt tgctagtgaa agataagacc ttgcatagag caccttacta cttcctgttg 180
qatctttqct qttcaqatat cctcaqatct gcaatttgtt tcccatttgt gttcaactct 240
qtcaaaaatq qctctacctq qacttatqqq actctgactt gcaaaqtgat tgcctttctg 300
ggggttttgt cctgtttcca cactgctttc atgctcttct gcatcagtgt caccagatac 360
ttagctatcg cccatcaccg cttctataca aagaggctga ccttttggac gtgtctggct 420
qtqatctqta tqqtqtqqac tctqtctqtq qccatqqcat ttcccccqqt tttaqacqtq 480
ggcacttact cattcattag ggaggaagat caatgcacct tccaacaccg ctccttcagg 540
gctaatgatt ccttaggatt tatgctgctt cttgctctca tcctcctagc cacacagctt 600
gtctacctca agctgatatt tttcgtccac gatcgaagaa aaatgaagcc agtccagttt 660
qtaqcaqcaq tcaqccaqaa ctqgactttt catqgtcctg gagccagtgg ccaggcagct 720
qccaattggc tagcaggatt tggaaggggt cccacaccac ccaccttgct gggcatcagg 780
caaaatgcaa acaccacagg cagaagaagg ctattggtct tagacgagtt caaaatggag 840
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tacciggtgg cctgttattg gagagttttt gcaagagggc ctgtagtacc agggggattt 960
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<210> 56
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<211> 364

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<213> Homo sapiens
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Ser Leu Pro Thr Met Gly Ala Asn Val Ser Gln Asp Asn Gly Thr Gly
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His Asn Ala Thr Phe Ser Glu Pro Leu Pro Phe Leu Tyr Val Leu Leu
                           40
Pro Ala Val Tyr Ser Gly Ile Cys Ala Val Gly Leu Thr Gly Asn Thr
                       55
                                           60
Ala Val Ile Leu Val Ile Leu Arg Ala Pro Lys Met Lys Thr Val Thr
                   70
Asn Val Phe Ile Leu Asn Leu Ala Val Ala Asp Gly Leu Phe Thr Leu
               85
                                   90
Val Leu Pro Val Asn Ile Ala Glu His Leu Leu Gln Tyr Trp Pro Phe
                               105
Gly Glu Leu Leu Cys Lys Leu Val Leu Ala Val Asp His Tyr Asn Ile
                           120
Phe Ser Ser Ile Tyr Phe Leu Ala Val Met Ser Val Asp Arg Tyr Leu
                       135
                                           140
Val Val Leu Ala Thr Val Arg Ser Arg His Met Pro Trp Arg Thr Tyr
                   150
                                       155
Arg Gly Ala Lys Val Ala Ser Leu Cys Val Trp Leu Gly Val Thr Val
                                   170
Leu Val Leu Pro Phe Phe Ser Phe Ala Gly Val Tyr Ser Asn Glu Leu
                               185
Gln Val Pro Ser Cys Gly Leu Ser Phe Pro Trp Pro Glu Arg Val Trp
                           200
Phe Lys Ala Ser Arg Val Tyr Thr Leu Val Leu Gly Phe Val Leu Pro
                       215
Val Cys Thr Ile Cys Val Leu Tyr Thr Asp Leu Leu Arg Arg Leu Arg
                   230
                                       235
Ala Val Arg Leu Arg Ser Gly Ala Lys Ala Leu Gly Lys Ala Arg Arg
                                   250
Lys Val Thr Val Leu Val Leu Val Leu Ala Val Cys Leu Leu Cys
                               265
Trp Thr Pro Phe His Leu Ala Ser Val Val Ala Leu Thr Thr Asp Leu
                           280
Pro Gln Thr Pro Leu Val Ile Ser Met Ser Tyr Val Ile Thr Ser Leu
                       295
Ser Tyr Ala Asn Ser Cys Leu Asn Pro Phe Leu Tyr Ala Phe Leu Asp
                   310
                                       315
Asp Asn Phe Arg Lys Asn Phe Arg Ser Ile Leu Arg Cys Ala Ala Ala
               325
                                   330
Arg Gly Arg Thr Pro Pro Ser Leu Gly Pro Gln Asp Glu Ser Cys Thr
                               345
Thr Ala Ser Ser Ser Leu Ala Lys Asp Thr Ser Ser
<210> 57
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<212> PRT

<212> DNA

<213> Homo sapiens

<400> 57

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atgggtgcca acgtctctca ggacaatggc actggccaca atgccacctt ctccgagcca 120

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ctgccgttcc tctatgtgct cctgcccgcc gtgtactccg ggatctgtgc tgtggggctg 180
actggcaaca cggccgtcat ccttgtaatc ctaagggcgc ccaagatgaa gacggtgacc 240
aacqtqttca teetgaacet ggeegtegee gaegggetet teaegetggt aetgeeegte 300
aacatcgcgg agcacctgct gcagtactgg cccttcgggg agctgctctg caagctggtg 360
ctggccgtcg accactacaa catcttctcc agcatctact tcctagccgt gatgagcgtg 420
gaccqatacc tggtggtgct ggccaccgtg aggtcccgcc acatgccctg gcgcacctac 480
cqqqqqqqa aqqtcqccaq cctqtqtqtc tqqctqqqcq tcacqqtcct qqttctqccc 540
ttettetett tegetggegt etacageaac gagetgeagg teceaagetg tgggetgage 600
ttcccgtggc ccgagcgggt ctggttcaag gccagccgtg tctacacttt ggtcctgggc 660
ttcgtgctgc ccgtgtgcac catctgtgtg ctctacacag acctcctgcg caggctgcgg 720
gccgtgcggc tccgctctgg agccaaggct ctaggcaagg ccaggcggaa ggtgaccgtc 780
ctggtcctcg tcgtgctggc cgtgtgcctc ctctgctgga cgcccttcca cctggcctct 840
gtcgtqgccc tgaccacgga cctgccccag accccactgg tcatcagtat gtcctacgtc 900
atcaccagee teagetacge caactegtge etgaacceet teetetacge etttetagat 960
gacaacttcc ggaagaactt ccgcagcata ttgcggtgcg cggccgcacg gggacgcacc 1020
ccacccagce tgggtcccca agatgagtcc tgcaccaccg ccagctcctc cctggccaag 1080
gacacttcat cqtqa
<210> 58
<211> 419
<212> PRT
<213> Homo sapiens
<400> 58
Met Cys Phe Ser Pro Ile Leu Glu Ile Asn Met Gln Ser Glu Ser Asn
Ile Thr Val Arg Asp Asp Ile Asp Asp Ile Asn Thr Asn Met Tyr Gln
Pro Leu Ser Tyr Pro Leu Ser Phe Gln Val Ser Leu Thr Gly Phe Leu
                            40
Met Leu Glu Ile Val Leu Gly Leu Gly Ser Asn Leu Thr Val Leu Val
Leu Tyr Cys Met Lys Ser Asn Leu Ile Asn Ser Val Ser Asn Ile Ile
                    70
Thr Met Asn Leu His Val Leu Asp Val Ile Ile Cys Val Gly Cys Ile
Pro Leu Thr Ile Val Ile Leu Leu Ser Leu Glu Ser Asn Thr Ala
                                105
Leu Ile Cys Cys Phe His Glu Ala Cys Val Ser Phe Ala Ser Val Ser
        115
                            120
Thr Ala Ile Asn Val Phe Ala Ile Thr Leu Asp Arg Tyr Asp Ile Ser
                        135
Val Lys Pro Ala Asn Arg Ile Leu Thr Met Gly Arg Ala Val Met Leu
                    150
                                        155
Met Ile Ser Ile Trp Ile Phe Ser Phe Phe Ser Phe Leu Ile Pro Phe
                165
                                    170
Ile Glu Val Asn Phe Phe Ser Leu Gln Ser Gly Asn Thr Trp Glu Asn
                                185
            180
Lys Thr Leu Leu Cys Val Ser Thr Asn Glu Tyr Tyr Thr Glu Leu Gly
                            200
                                                205
        195
Met Tyr Tyr His Leu Leu Val Gln Ile Pro Ile Phe Phe Phe Thr Val
                        215
                                            220
    210
Val Val Met Leu Ile Thr Tyr Thr Lys Ile Leu Gln Ala Leu Asn Ile
                                        235
                    230
Arg Ile Gly Thr Arg Phe Ser Thr Gly Gln Lys Lys Lys Ala Arg Lys
                                    250
                245
Lys Lys Thr Ile Ser Leu Thr Thr Gln His Glu Ala Thr Asp Met Ser
                                265
            260
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Gln Ser Ser Gly Gly Arg Asn Val Val Phe Gly Val Arg Thr Ser Val
                            280
Ser Val Ile Ile Ala Leu Arg Arg Ala Val Lys Arg His Arg Glu Arg
                                            300
                        295
Arg Glu Arg Gln Lys Arg Val Phe Arg Met Ser Leu Leu Ile Ile Ser
                    310
                                        315
Thr Phe Leu Leu Cys Trp Thr Pro Ile Ser Val Leu Asn Thr Thr Ile
                325
                                    330
Leu Cys Leu Gly Pro Ser Asp Leu Leu Val Lys Leu Arg Leu Cys Phe
                                                    350
            340
                                345
Leu Val Met Ala Tyr Gly Thr Thr Ile Phe His Pro Leu Leu Tyr Ala
                            360
                                                365
Phe Thr Arg Gln Lys Phe Gln Lys Val Leu Lys Ser Lys Met Lys Lys
                        375
                                            380
Arg Val Val Cys Ala Ala Ala Arg Gly Arg Thr Pro Pro Ser Leu Gly
                    390
                                        395
Pro Gln Asp Glu Ser Cys Thr Thr Ala Ser Ser Ser Leu Ala Lys Asp
                                    410
Thr Ser Ser
<210> 59
<211> 1260
<212> DNA
<213> Homo sapiens
<400> 59
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gatgacattg atgacatcaa caccaatatg taccaaccac tatcatatcc gttaagcttt 120
caagtgtete teaceggatt tettatgtta gaaattgtgt tgggaettgg cageaacete 180
actgtattgg tactttactg catgaaatcc aacttaatca actctgtcag taacattatt 240
acaatgaatc ttcatgtact tgatgtaata atttgtgtgg gatgtattcc tctaactata 300
qttalcotto tgctttcact ggagagtaac actgctctca tttgctgttt ccatgaggct 360
tgtgtatctt ttgcaagtgt ctcaacagca atcaacgttt ttgctatcac tttggacaga 420
tatqacatct ctqtaaaacc tqcaaaccqa attctqacaa tqqqcaqaqc tqtaatqtta 480
atgatatcca tttggatttt ttctttttc tctttcctga ttccttttat tgaggtaaat 540
tttttcagtc ttcaaagtgg aaatacctgg gaaaacaaga cacttttatg tgtcagtaca 600
aatgaatact acactgaact gggaatgtat tatcacctgt tagtacagat cccaatattc 660
tttttcactg ttgtagtaat gttaatcaca tacaccaaaa tacttcaggc tcttaatatt 720
cgaataggca caagattttc aacagggcag aagaagaaag caagaaagaa aaagacaatt 780
tototaacca cacaacatga ggotacagac atgtoacaaa gcagtggtgg gagaaatgta 840
gtctttggtg taagaacttc agtttctgta ataattgccc tccggcgagc tgtgaaacga 900
caccgtgaac gacgagaaag acaaaagaga gtcttcagga tgtctttatt gattatttct 960
acatttette tetgetggae accaatttet gttttaaata ceaccatttt atgtttagge 1020
ccaaqtgacc ttttagtaaa attaagattg tgttttttag tcatggctta tggaacaact 1080
atatttcacc ctctattata tgcattcact agacaaaaat ttcaaaaaggt cttgaaaagt 1140
aaaatgaaaa agcgagttgt ttgtgcggcc gcacggggac gcaccccacc cagcctgggt 1200
ccccaaqatq aqtcctgcac caccqccagc tectecetgg ccaaggacac ttcategtga 1260
<210> 60
<211> 370
<212> PRT
<213> Homo sapiens
<400> 60
Met Gly Gln Pro Gly Asn Gly Ser Ala Phe Leu Leu Ala Pro Asn Arg
                                    10
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Ser His Ala Pro Asp His Asp Val Thr Gln Gln Arg Asp Glu Val Trp
                               25
Val Val Gly Met Gly Ile Val Met Ser Leu Ile Val Leu Ala Ile Val
                           40
Phe Gly Asn Val Leu Val Ile Thr Ala Ile Ala Lys Phe Glu Arg Leu
                       55
Gln Thr Val Thr Asn Tyr Phe Ile Thr Ser Leu Ala Cys Ala Asp Leu
                  70
Val Met Gly Leu Ala Val Val Pro Phe Gly Ala Ala His Ile Leu Met
                                  90
Lys Met Trp Thr Phe Gly Asn Phe Trp Cys Glu Phe Trp Thr Ser Ile
          100
                              105
Asp Val Leu Cys Val Thr Ala Ser Ile Glu Thr Leu Cys Val Ile Ala
                          120
 . 115
Val Asp Arg Tyr Phe Ala Ile Thr Ser Pro Phe Lys Tyr Gln Ser Leu
                      135
                                          140
Leu Thr Lys Asn Lys Ala Arg Val Ile Ile Leu Met Val Trp Ile Val
                  150
                                      155
Ser Gly Leu Thr Ser Phe Leu Pro Ile Gln Met His Trp Tyr Arg Ala
              165
                                  170
Thr His Gln Glu Ala Ile Asn Cys Tyr Ala Asn Glu Thr Cys Cys Asp
                              185
Phe Phe Thr Asn Gln Ala Tyr Ala Ile Ala Ser Ser Ile Val Ser Phe
                          200
Tyr Val Pro Leu Val Ile Met Val Phe Val Tyr Ser Arg Val Phe Gln
                       215
Glu Ala Lys Arg Gln Leu Gln Lys Ile Asp Lys Ser Glu Gly Arg Phe
                                      235
His Val Gln Asn Leu Ser Gln Val Glu Gln Asp Gly Arg Thr Gly His
                                   250
Gly Leu Arg Arg Ser Ser Lys Phe Cys Leu Lys Glu His Lys Ala Leu
                               265
Lys Thr Leu Gly Ile Ile Met Gly Thr Phe Thr Leu Cys Trp Leu Pro
                           280.
Phe Phe Ile Val Asn Ile Val His Val Ile Gln Asp Asn Leu Ile Arg
                       295
Lys Glu Val Tyr Ile Leu Leu Asn Trp Ile Gly Tyr Val Asn Ser Gly
                                       315
Phe Asn Pro Leu Ile Tyr Cys Arg Ser Pro Asp Phe Arg Ile Ala Phe
                                   330
Gln Glu Leu Cys Ala Arg Gly Arg Thr Pro Pro Ser Leu Gly Pro
                              345
Gln Asp Glu Ser Cys Thr Thr Ala Ser Ser Ser Leu Ala Lys Asp Thr
                           360
Ser Ser
  370
<210> 61
<211> 382
<212> PRT
<213> Homo sapiens
<400> 61
Met Asp Ser Ser Thr Gly Pro Gly Asn Thr Ser Asp Cys Ser Asp Pro
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Leu Ala Gln Ala Ser Cys Ser Pro Ala Pro Gly Ser Trp Leu Asn Leu
           20
                               25
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Ser His Val Asp Gly Asn Gln Ser Asp Pro Cys Gly Leu Asn Arg Thr

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35
                         40
                                            45
Gly Leu Gly Gly Asn Asp Ser Leu Cys Pro Gln Thr Gly Ser Pro Ser
                    55
Met Val Thr Ala Ile Thr Ile Met Ala Leu Tyr Ser Ile Val Cys Val
                              75
                 70
Val Gly Leu Phe Gly Asn Phe Leu Val Met Tyr Val Ile Val Arg Tyr
                                90
Thr Lys Met Lys Thr Ala Thr Asn Ile Tyr Ile Phe Asn Leu Ala Leu
                            105
Ala Asp Ala Leu Ala Thr Ser Thr Leu Pro Phe Gln Ser Val Asn Tyr
                         120
Leu Met Gly Thr Trp Pro Phe Gly Thr Ile Leu Cys Lys Ile Val Ile
                     135
                                 140
Ser Ile Asp Tyr Tyr Asn Met Phe Thr Ser Ile Phe Thr Leu Cys Thr
            150 155
Met Ser Val Asp Arg Tyr Ile Ala Val Cys His Pro Val Lys Ala Leu
             165 170 175
Asp Phe Arg Thr Pro Arg Asn Ala Lys Ile Val Asn Val Cys Asn Trp
                            185
Ile Leu Ser Ser Ala Ile Gly Leu Pro Val Met Phe Met Ala Thr Thr
                         200
Lys Tyr Arg Gln Gly Ser Ile Asp Cys Thr Leu Thr Phe Ser His Pro
                     215
Thr Trp Tyr Trp Glu Asn Leu Leu Lys Ile Cys Val Phe Ile Phe Ala
                 230
                                    235
Phe Ile Met Pro Ile Leu Ile Ile Thr Val Cys Tyr Gly Leu Met Ile
                                250
Leu Arg Leu Lys Ser Val Arg Met Leu Ser Gly Ser Lys Glu Lys Asp
                             265
Arg Asn Leu Arg Arg Ile Thr Arg Met Val Leu Val Val Val Ala Val
                         280
Phe Ile Val Cys Trp Thr Pro Ile His Ile Tyr Val Ile Ile Lys Ala
                      295
Leu Ile Thr Ile Pro Glu Thr Thr Phe Gln Thr Val Ser Trp His Phe
                  310
                                     315
Cys Ile Ala Leu Gly Tyr Thr Asn Ser Cys Leu Asn Pro Val Leu Tyr
              325
                                330
Ala Phe Leu Asp Glu Asn Phe Lys Arg Cys Phe Arg Glu Phe Cys Ala
                             345
Ala Ala Arg Gly Arg Thr Pro Pro Ser Leu Gly Pro Gln Asp Glu Ser
                         360
Cys Thr Thr Ala Ser Ser Ser Leu Ala Lys Asp Thr Ser Ser
                      375
<210> 62
<211> 382
<212> PRT
<213> Homo sapiens
<400> 62
Met Ala Pro Asn Thr Ser Thr Met Asp Glu Ala Gly Leu Pro Ala Glu
                                 10
Arg Asp Phe Ser Phe Arg Ile Leu Thr Ala Cys Phe Leu Ser Leu Leu
                              25
Ile Leu Ser Thr Leu Leu Gly Asn Thr Leu Val Cys Ala Ala Val Ile
                         40
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Arg Phe Arg His Leu Arg Ser Lys Val Thr Asn Phe Phe Val Ile Ser

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Leu Ala Val Ser Asp Leu Leu Val Ala Val Leu Val Met Pro Trp Lys
Ala Val Ala Glu Ile Ala Gly Phe Trp Pro Phe Gly Ser Phe Cys Asn
                                   90
Ile Trp Val Ala Phe Asp Ile Met Cys Ser Thr Ala Ser Ile Leu Asn
                               105
Leu Cys Val Ile Ser Val Asp Arg Tyr Trp Ala Ile Ser Ser Pro Phe
                           120
Gln Tyr Glu Arg Lys Met Thr Pro Lys Ala Ala Phe Ile Leu Ile Ser
                       135
Val Ala Trp Thr Leu Ser Val Leu Ile Ser Phe Ile Pro Val Gln Leu
                                       155
Ser Trp His Lys Ala Lys Pro Thr Trp Pro Leu Asp Gly Asn Phe Thr
                                   170
Ser Leu Glu Asp Thr Glu Asp Asp Asn Cys Asp Thr Arg Leu Ser Arg
                              185
Thr Tyr Ala Ile Ser Ser Ser Leu Ile Ser Phe Tyr Ile Pro Val Ala
                           200
Ile Met Ile Val Thr Tyr Thr Ser Ile Tyr Arg Ile Ala Gln Lys Gln
                       215
Ile Arq Arq Ile Ser Ala Leu Glu Arg Ala Ala Val His Ala Lys Asn
                   230
                                       235
Cys Gln Thr Thr Ala Gly Asn Gly Asn Pro Val Glu Cys Ala Gln Ser
               245
                                   250
Glu Ser Ser Phe Lys Met Ser Phe Lys Arg Glu Thr Lys Val Leu Lys
                               265
Thr Leu Ser Val Ile Met Gly Val Phe Val Cys Cys Trp Leu Pro Phe
                           280
Phe Ile Ser Asn Cys Met Val Pro Phe Cys Gly Ser Glu Glu Thr Gln
                       295
                                           300
Pro Phe Cys Ile Asp Ser Ile Thr Phe Asp Val Phe Val Trp Phe Gly
                                       315
                   310
Trp Ala Asn Ser Ser Leu Asn Pro Ile Ile Tyr Ala Phe Asn Ala Asp
                                   330
                325
Phe Gln Lys Ala Phe Ser Thr Leu Leu Gly Cys Tyr Arg Leu Cys Ala
                               345
            340
Ala Ala Arg Gly Arg Thr Pro Pro Ser Leu Gly Pro Gln Asp Glu Ser
       355
                           360
Cys Thr Thr Ala Ser Ser Ser Leu Ala Lys Asp Thr Ser Ser
                        375
    370
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<210> 63
<211> 451
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<400> 63

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        Val
        Leu
        Ser
        Pro
        Gly
        Gly
        Asn
        Asn
        Thr
        Gly
        Ile
        Ser
        Asp
        Val
        Thr

        Val
        Ser
        Tyr
        Gln
        Val
        Ile
        Thr
        Ser
        Leu
        Leu
        Leu
        Gly
        Thr
        Leu
        Ile
        Pro
        Pro
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<212> PRT

<213> Homo sapiens

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Gln Val Leu Asn Lys Trp Thr Leu Gly Gln Val Thr Cys Asp Leu Phe
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          100
                                               110
Ile Ala Leu Asp Val Leu Cys Cys Thr Ser Ser Ile Leu His Leu Cys
                                      125
                    120
Ala Ile Ala Leu Asp Arg Tyr Trp Ala Ile Thr Asp Pro Ile Asp Tyr
                      135
                               140
Val Asn Lys Arg Thr Pro Arg Arg Ala Ala Ala Leu Ile Ser Leu Thr
                  150
                                     155
Trp Leu Ile Gly Phe Leu Ile Ser Ile Pro Pro Met Leu Gly Trp Arg
              165
                                 170
Thr Pro Glu Asp Arg Ser Asp Pro Asp Ala Cys Thr Ile Ser Lys Asp
                              185
His Gly Tyr Thr Ile Tyr Ser Thr Phe Gly Ala Phe Tyr Ile Pro Leu
                          200
                                      205
Leu Leu Met Leu Val Leu Tyr Gly Arg Ile Phe Arg Ala Ala Arg Phe
                      215
                                         220
Arg Ile Arg Lys Thr Val Lys Lys Val Glu Lys Thr Gly Ala Asp Thr
                  230
                                      235
Arg His Gly Ala Ser Pro Ala Pro Gln Pro Lys Lys Ser Val Asn Gly
               245
                                  250
Glu Ser Gly Ser Arg Asn Trp Arg Leu Gly Val Glu Ser Lys Ala Gly
                              265
Gly Ala Leu Cys Ala Asn Gly Ala Val Arg Gln Gly Asp Asp Gly Ala
                          280
Ala Leu Glu Val Ile Glu Val His Arg Val Gly Asn Ser Lys Glu His
                      295
Leu Pro Leu Pro Ser Glu Ala Gly Pro Thr Pro Cys Ala Pro Ala Ser
                   310
                                      315
Phe Glu Arg Lys Asn Glu Arg Asn Ala Glu Ala Lys Arg Lys Met Ala
               325
                                  330
Leu Ala Arg Glu Arg Lys Thr Val Lys Thr Leu Gly Ile Ile Met Gly
                              345
           340
Thr Phe Ile Leu Cys Trp Leu Pro Phe Phe Ile Val Ala Leu Val Leu
                           360
       355
Pro Phe Cys Glu Ser Ser Cys His Met Pro Thr Leu Leu Gly Ala Ile
                       375
Ile Asn Trp Leu Gly Tyr Ser Asn Ser Leu Leu Asn Pro Val Ile Tyr
                  390
                                      395
Ala Tyr Phe Asn Lys Asp Phe Gln Asn Ala Phe Lys Lys Ile Ile Lys
               405
                                  410
Cys Asn Phe Cys Ala Ala Ala Arg Gly Arg Thr Pro Pro Ser Leu Gly
                              425
                                                 430
          420
Pro Gln Asp Glu Ser Cys Thr Thr Ala Ser Ser Leu Ala Lys Asp
                           440
       435
Thr Ser Ser
  450
<210> 64
<211> 394
<212> PRT
<213> Homo sapiens
<400> 64
Met Ala Pro Trp Pro His Glu Asn Ser Ser Leu Ala Pro Trp Pro Asp
                - 5
                                  10
Leu Pro Thr Leu Ala Pro Asn Thr Ala Asn Thr Ser Gly Leu Pro Gly
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Val Pro Trp Glu Ala Ala Leu Ala Gly Ala Leu Leu Ala Leu Ala Val
Leu Ala Thr Val Gly Gly Asn Leu Leu Val Ile Val Ala Ile Ala Trp
Thr Pro Arg Leu Gln Thr Met Thr Asn Val Phe Val Thr Ser Leu Ala
Ala Ala Asp Leu Val Met Gly Leu Leu Val Val Pro Pro Ala Ala Thr
Leu Ala Leu Thr Gly His Trp Pro Leu Gly Ala Thr Gly Cys Glu Leu
                               105
Trp Thr Ser Val Asp Val Leu Cys Val Thr Ala Ser Ile Glu Thr Leu
                           120
Cys Ala Leu Ala Val Asp Arg Tyr Leu Ala Val Thr Asn Pro Leu Arg
                       135
Tyr Gly Ala Leu Val Thr Lys Arg Cys Ala Arg Thr Ala Val Val Leu
                   150
Val Trp Val Val Ser Ala Ala Val Ser Phe Ala Pro Ile Met Ser Gln
                                   170
               165
Trp Trp Arg Val Gly Ala Asp Ala Glu Ala Gln Arg Cys His Ser Asn
                               185
Pro Arg Cys Cys Ala Phe Ala Ser Asn Met Pro Tyr Val Leu Leu Ser
                           200
Ser Ser Val Ser Phe Tyr Leu Pro Leu Leu Val Met Leu Phe Val Tyr
                       215
Ala Arg Val Phe Val Val Ala Thr Arg Gln Leu Arg Leu Leu Arg Gly
                   230
                                        235
Glu Leu Gly Arg Phe Pro Pro Glu Glu Ser Pro Pro Ala Pro Ser Arg
                                   250
                245
Ser Leu Ala Pro Ala Pro Val Gly Thr Cys Ala Pro Pro Glu Gly Val
                               265
Pro Ala Cys Gly Arg Arg Pro Ala Arg Leu Leu Pro Leu Arg Glu His
                            280
Arg Ala Leu Cys Thr Leu Gly Leu Ile Met Gly Thr Phe Thr Leu Cys
                                            300
                        295
Trp Leu Pro Phe Phe Leu Ala Asn Val Leu Arg Ala Leu Gly Gly Pro
                                       315
                   310
Ser Leu Val Pro Gly Pro Ala Phe Leu Ala Leu Asn Trp Leu Gly Tyr
               325
                                    330
Ala Asn Ser Ala Phe Asn Pro Leu Ile Tyr Cys Arg Ser Pro Asp Phe
                               345
            340
Arg Ser Ala Phe Arg Arg Leu Leu Cys Arg Cys Ala Ala Ala Arg Gly
                           360
Arg Thr Pro Pro Ser Leu Gly Pro Gln Asp Glu Ser Cys Thr Thr Ala
                        375
Ser Ser Ser Leu Ala Lys Asp Thr Ser Ser
                    390
<210> 65
<211> 362
<212> PRT
<213> Homo sapiens
<400> 65
Met Gly Pro Thr Ser Val Pro Leu Val Lys Ala His Arg Ser Ser Val
                                    10
Ser Asp Tyr Val Asn Tyr Asp Ile Ile Val Arg His Tyr Asn Tyr Thr
                                25
            20
Gly Lys Leu Asn Ile Ser Ala Asp Lys Glu Asn Ser Ile Lys Leu Thr
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40
Ser Val Val Phe Ile Leu Ile Cys Cys Phe Ile Ile Leu Glu Asn Ile
Phe Val Leu Leu Thr Ile Trp Lys Thr Lys Lys Phe His Arg Pro Met
                 70
                                  75
Tyr Tyr Phe Ile Gly Asn Leu Ala Leu Ser Asp Leu Leu Ala Gly Val
                               90
Ala Tyr Thr Ala Asn Leu Leu Ser Gly Ala Thr Thr Tyr Lys Leu
    100
                           105
Thr Pro Ala Gln Trp Phe Leu Arg Glu Gly Ser Met Phe Val Ala Leu
 115
                     120
Ser Ala Ser Val Phe Ser Leu Leu Ala Ile Ala Ile Glu Arg Tyr Ile
                    135
Thr Met Leu Lys Met Lys Leu His Asn Gly Ser Asn Asn Phe Arg Leu
145 150
                                 155
Phe Leu Leu Ile Ser Ala Cys Trp Val Ile Ser Leu Ile Leu Gly Gly
                               170
             165
Leu Pro Ile Met Gly Trp Asn Cys Ile Ser Ala Leu Ser Ser Cys Ser
                           185
    180
Thr Val Leu Pro Leu Tyr His Lys His Tyr Ile Leu Phe Cys Thr Thr
 195
                        200
Val Phe Thr Leu Leu Leu Ser Ile Val Ile Leu Tyr Cys Arg Ile
                                      220
       215
Tyr Ser Leu Val Arg Thr Arg Ser Arg Arg Leu Thr Phe Arg Lys Asn
                                  235
                230
Ile Ser Lys Ala Ser Arg Ser Ser Glu Lys Ser Leu Ala Leu Leu Lys
                            250
       245
Thr Val Ile Ile Val Leu Ser Val Phe Ile Ala Cys Trp Ala Pro Leu
   260
                            265
Phe Ile Leu Leu Leu Leu Asp Val Gly Cys Lys Val Lys Thr Cys Asp
                                         285
                        280
       275
Ile Leu Phe Arg Ala Glu Tyr Phe Leu Val Leu Ala Val Leu Asn Ser
                                      300
                     295
Gly Thr Asn Pro Ile Ile Tyr Thr Leu Thr Asn Lys Glu Met Arg Arg
                 310
                                   315
Ala Phe Ile Arg Ile Met Ser Cys Cys Lys Cys Ala Ala Ala Arg Gly
                              330
                                     335
              325
Arg Thr Pro Pro Ser Leu Gly Pro Gln Asp Glu Ser Cys Thr Thr Ala
          340
                        345
Ser Ser Ser Leu Ala Lys Asp Thr Ser Ser
                        360
```